Figure S2: Mean precision values at 20% recall (P20R) within each evaluation category. (**a**) Evaluation of initial submissions using held-out genes. Biological Process (GO-BP), Cellular Component (GO-CC), and Molecular Function (GO-MF) branches are indicated on the x-axis, grouped by specificity (indicated by the minimum number of genes in the training set associated with each GO term in a given category). (**b**) Mean P20R within each evaluation category, evaluated prospectively using newly-annotated genes. (**c**) (**d**) As described in (a),(b), with colored region indicating performance of second-round submissions and grey region indicating performance in the first set of submissions, for comparison.